

FIGURE 1A

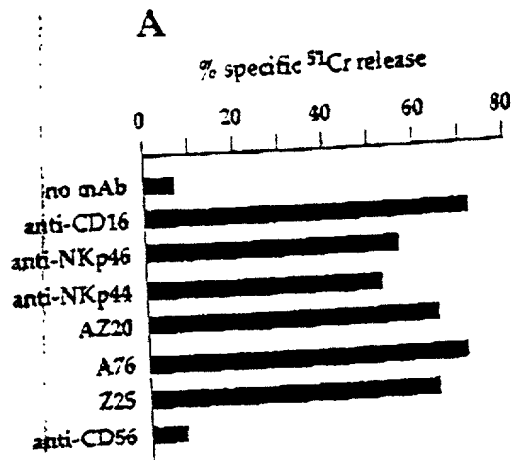


Fig 1A

FIGURE 1B

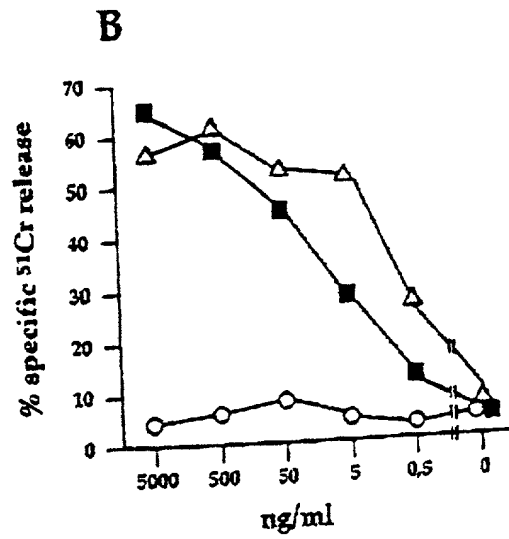


Fig 1B

FIGURE 1C

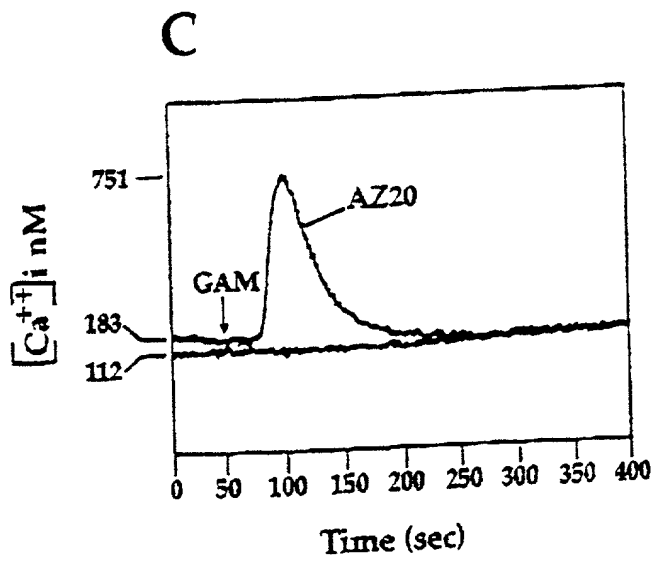


Fig 1C

FIGURE 2A

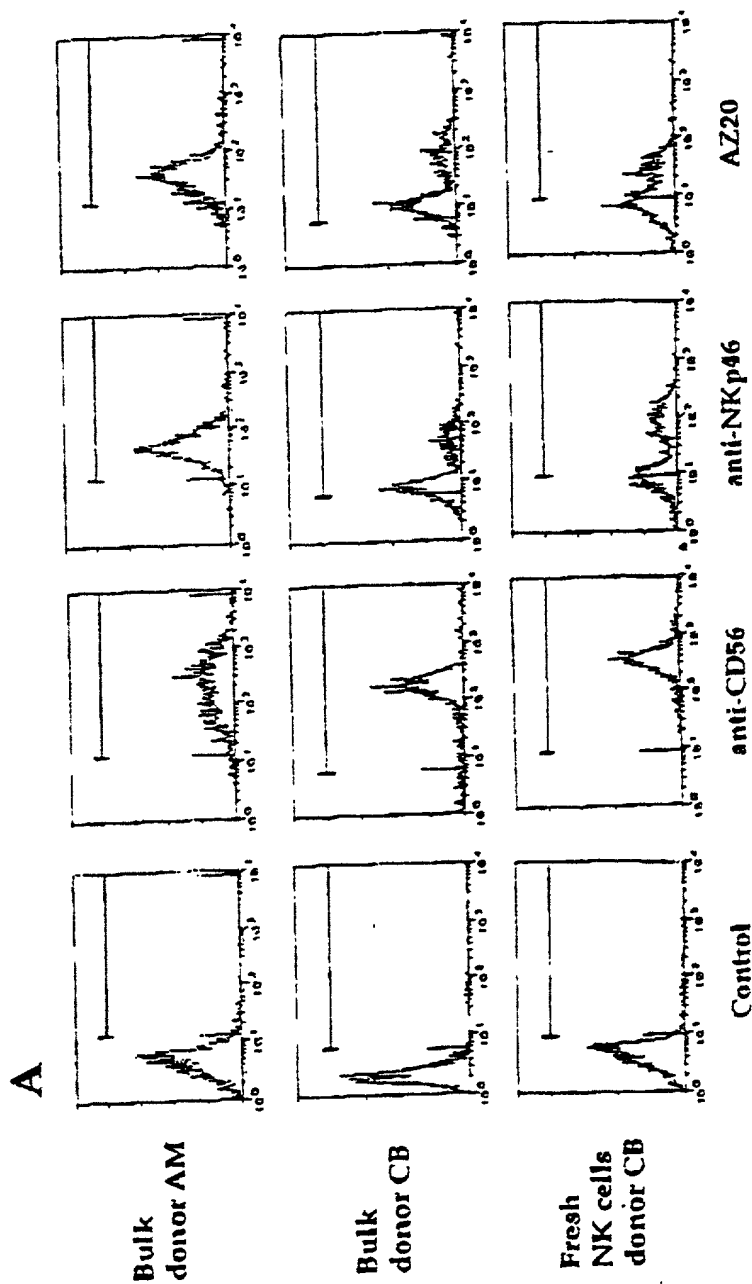


FIGURE 2B

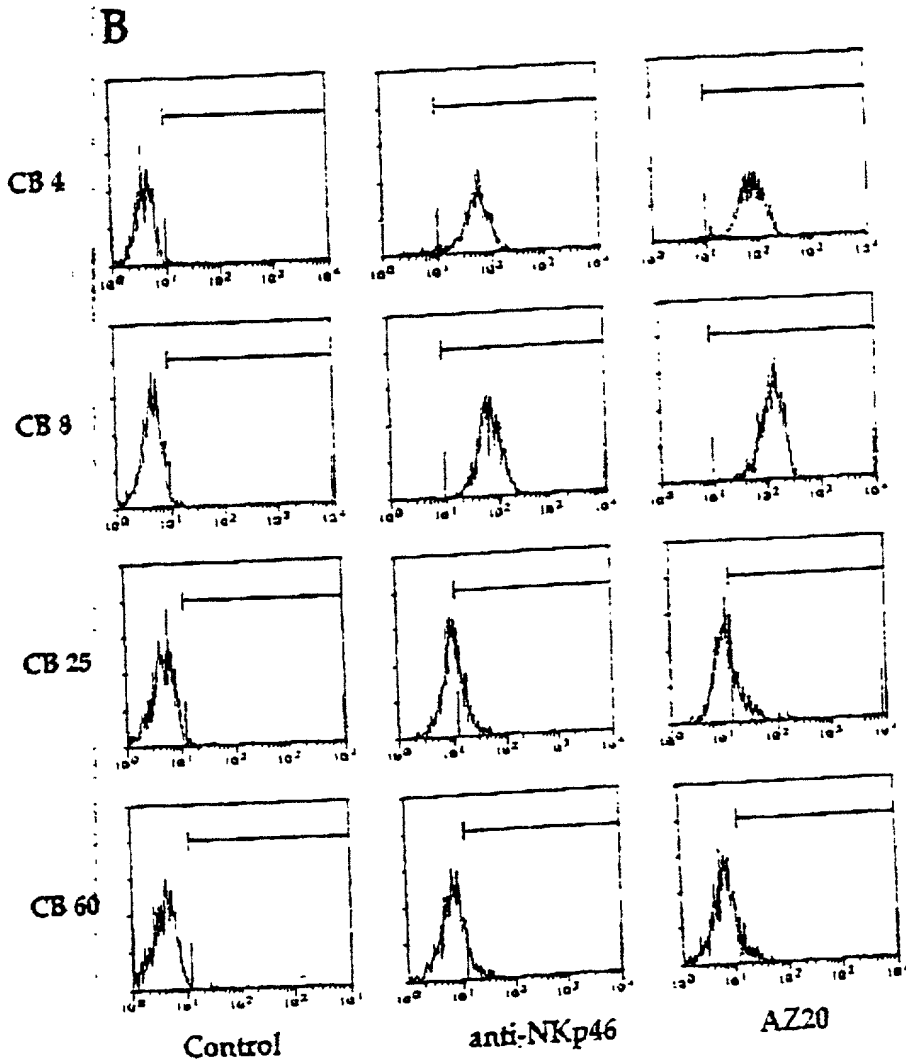


Fig 2B

7/19  
Drawing sheet

FIGURE 3B

**B**

Cells Daudi NK

69 —

45 —

28 —

18 —



Blot

AZ20

Fig 3B

FIGURE 4A

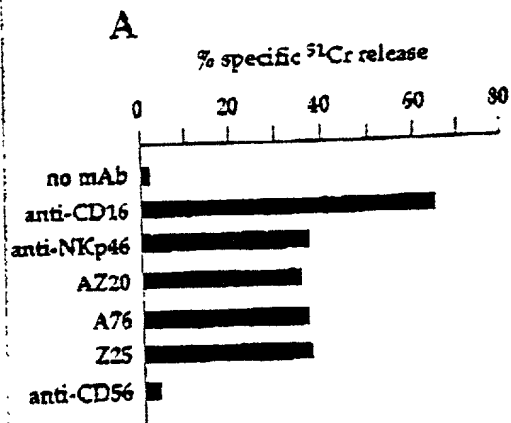


Fig 4A

FIGURE 3A

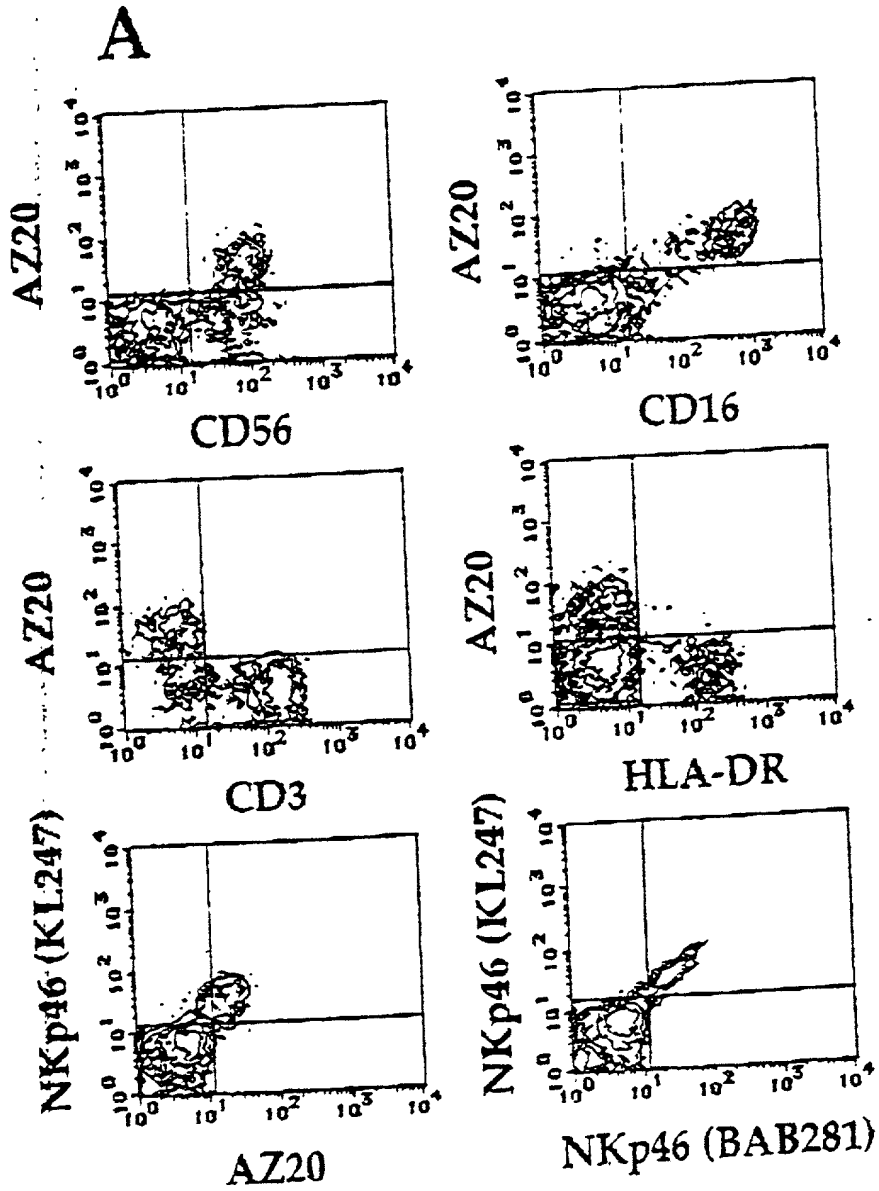




FIGURE 4B

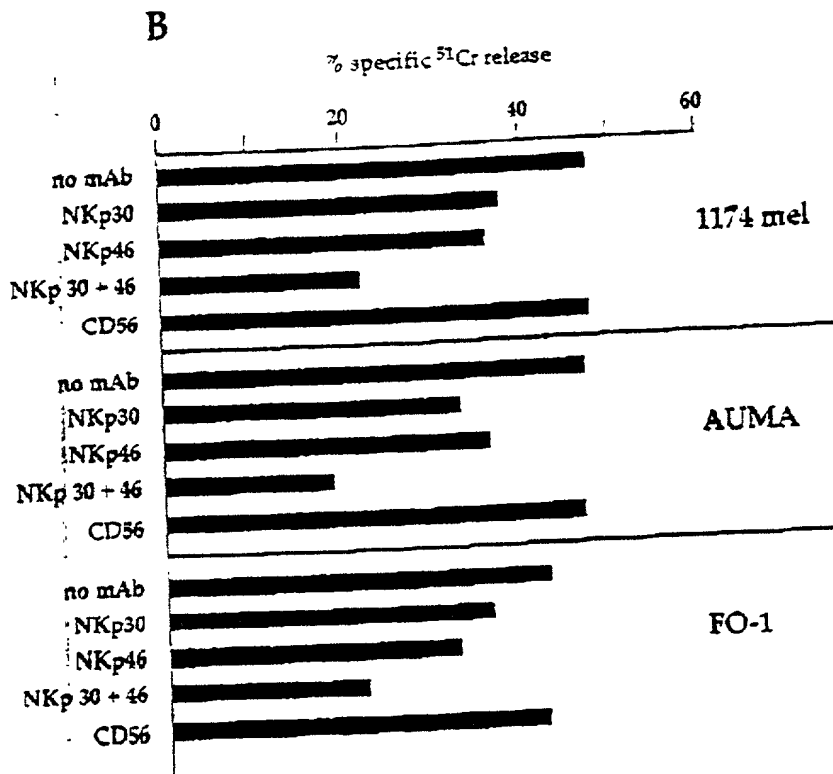


FIGURE 5

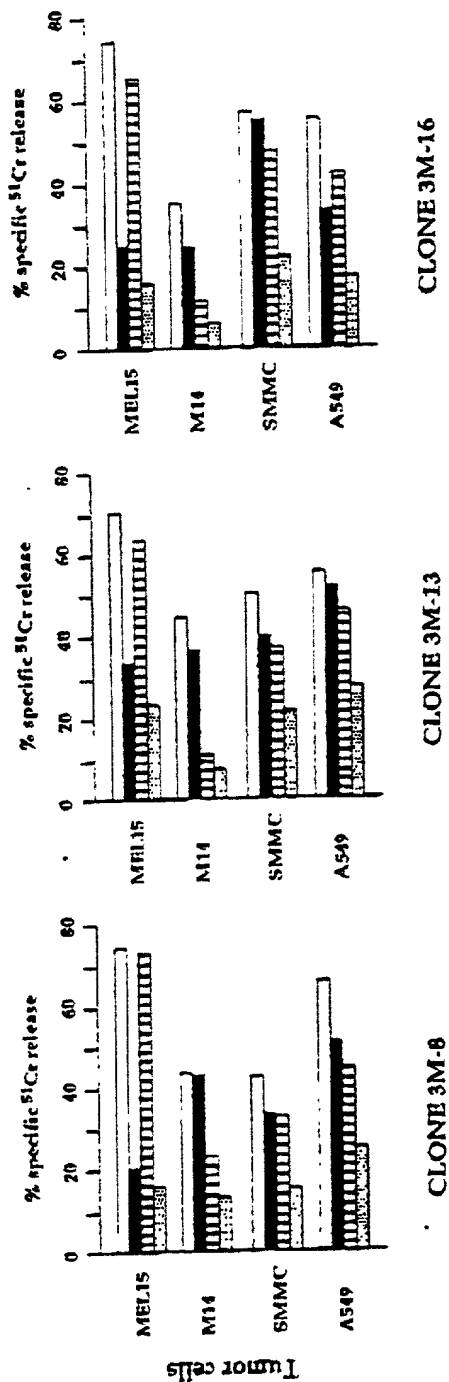


FIGURE 6A

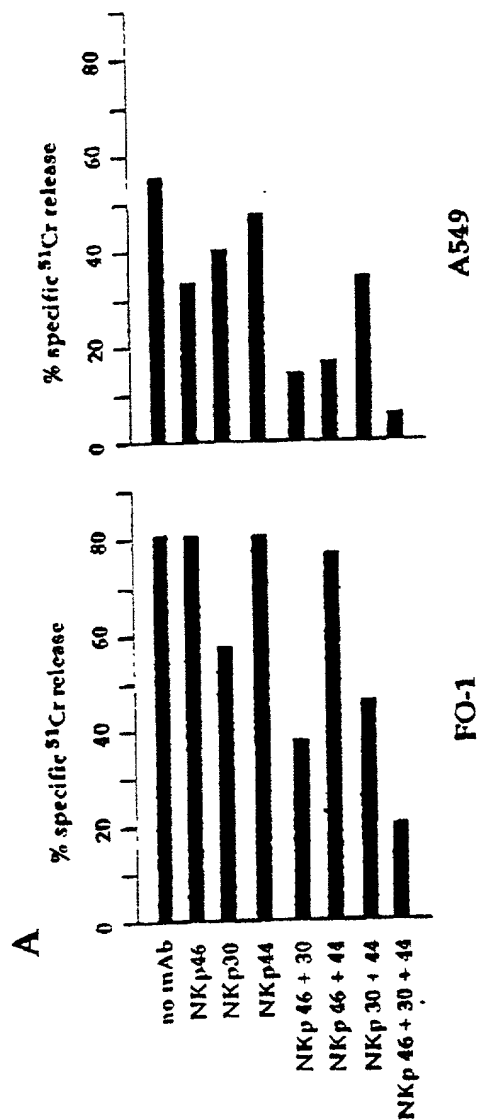
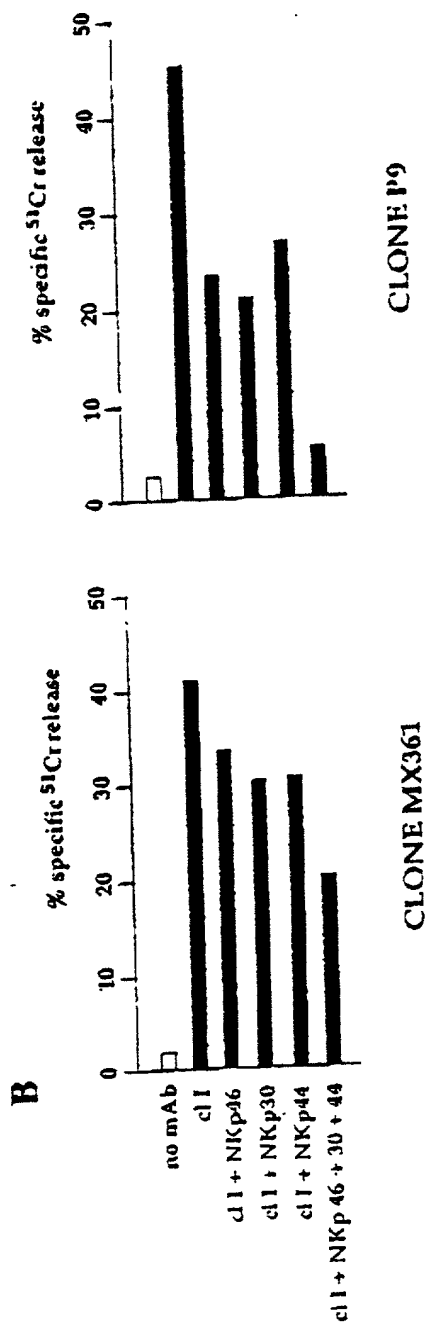


FIGURE 6B



F<sub>18</sub> 7A

FIGURE 7A

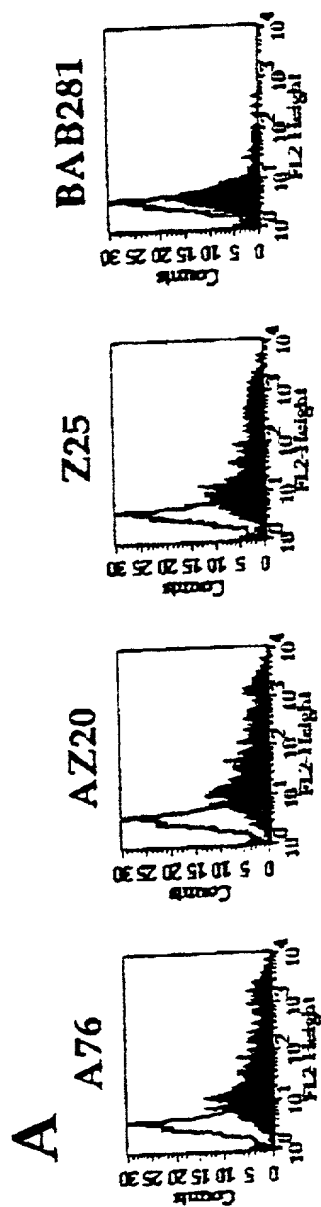
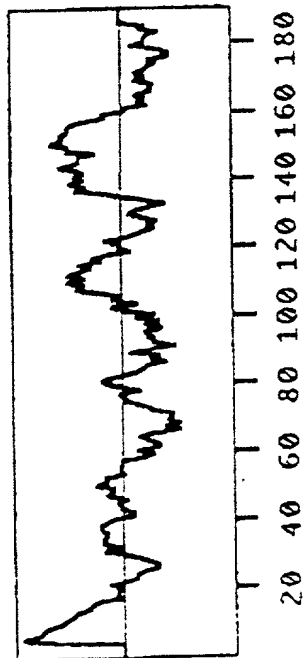


FIGURE 7B

**B** SEQUENCE N° 2

mowlllil	mvhpgscalw	VSQPPEIRTL	EGSSAFLPOS	<del>FN</del> ASQGLAI	50
GSVTWFERDEV	VPKKEVRNGT	PEFRGLAPL	ASSRFLHDHQ	AELHIRDVRG	100
HDASIYVCRV	EVLGLGVGTG	<del>NG</del> TRLVVEKE	HPQLGAGTVL	<u>LLRAGFYAVS</u>	150
<u>FLSVAVGSTV</u>	YYQKCHCHM	GTHCHSSDGP	RGVPEPRCP		190



15/19  
Drawing sheet

1 ccttcctcct ccacccagac ctcactgctc agatccccc cgcacaactgg gacatcttcc  
61 gacatggcct ggatgctgtt gctcatcttg atcatgggccc atccaggatc ctgtgctctc  
121 tgggtgtccc agccccctga gattcgtacc ctggaaggat cctctgcctt cctgcccctgc  
181 tccctcaatg ccagccaagg gagactggcc attggctccg tcacgtgggt ccgagatgag  
241 gtggttccag ggaaggaggt gaggaatgga accccagagt tcaggggccg cctggcccca  
301 cttgcttctt cccgtttcct ccattgaccac caggctgagc tgcacatccg ggcgtgcca  
361 ggccatgacg ccagcatcta cgtgtgcaga gtggagggtc tgggccttgg tgtcgggaca  
421 gggaaatggga ctgggtggt ggtggagaaa gaacatccct agctaggggc tggtagagtc  
481 ctctctcttc gggctggatt ctatgctgtc agctttctct ctgtggccgt gggcagcacc  
541 gtctattacc agggcaaatg ccactgtcac atgggaacac actgccactc cttagatggg  
601 ccccgaggcg tgattccaga gccagatgc cctagtccct cttcaaaaga ccccaataaa  
661 tctgccccac cact

SEQ ID N°1

FIGURE 7C

Fig 7C

F.88A

FIGURE 8A

A

1 2 3 4 5 6 7 8 9 10

28S—

18S—





FIGURE 8B

B

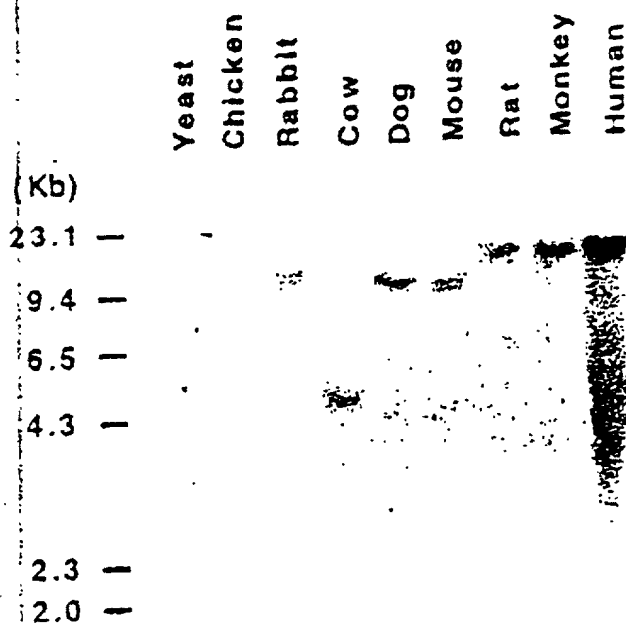


Fig 8B

FIGURE 9A

A

Cells Daudi NK

69 —

45 —

28 —

18 —



Blot

I

Fig 3A

FIGURE 9B

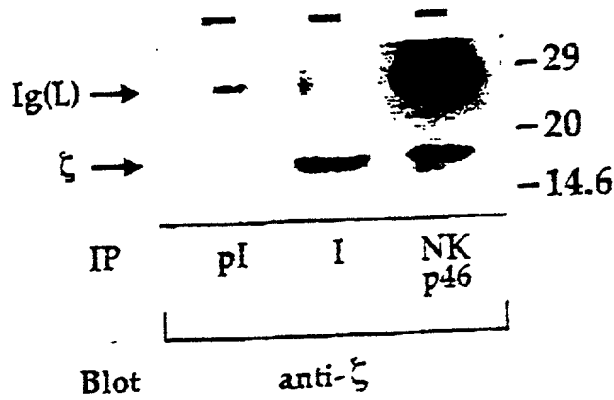
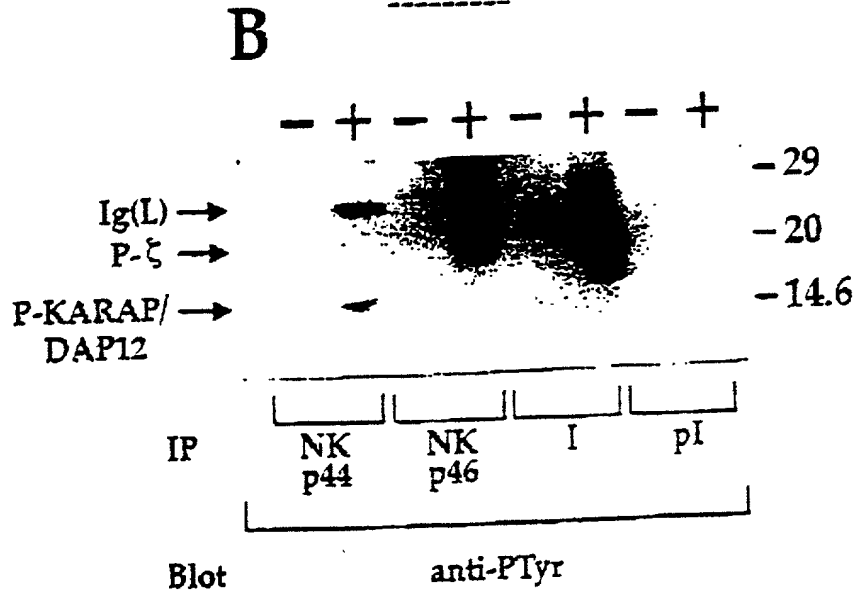


Fig 9B